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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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                                                                                                                                                                                                                                                                        seq length: 0 seq length: 2000000000
         Issued_Patents_AA:*
1: /cgn2_6/ptodata/2
2: /cgn2_6/ptodata/2
3: /cgn2_6/ptodata/2
4: /cgn2_6/ptodata/2
5: /cgn2_6/ptodata/2
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Gapop 10.0 , Gapext 0.5
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1713
1 MDIVVSGSHPLWVDSF
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Copyright (c) 1993 - 2000 Com
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/cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
/cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/2/iaa/B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/backfiles1.pep:*
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612.969 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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US-U9-1/2-9//-4		-08-555-568E	-977	-065-474-2	29A-	-485-	-196	-387-	9-196-387-	T-US95-08069	-09-519-223-2	899	-735-716	US-08-422-106-2	-281-193-	-474-	-810-71	-474-	-429A	œ	-313-3	-313-	13 - 3	-313-3	US-09-439-313-532	US-09-439-313-376	ID
Sequence 4, Appli	21,	e 21,	ω ·	23		23	e 2	10,					2, Appl	2, Appl	N	139, Ar	10, App	ω		ω Δ	377,	Sequence 378, App	380,	379,	e 532,	Sequence 376, App	Description

4.5	44	43	42	41	40	. 39	38	37	36	ن ان ان	34	ω ω	32	31	30	200	28
193.5	193.5	193.5	194.5	194.5	194.5	196	197.5	197.5	197.5	199	199	199.5	200.5	200.5	200.5	107	201
11.3	11.3	11.3	11.4	11.4	11.4	11.4	11.5	11.5	11.5	11.6	11.6	11.6	11.7	11.7	11.7	11.7	11.7
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US-08-185-432-18	US-09-113-825-3	US-08-537-210A-3	US-09-065-474-28	US-08-847-429A-28	US-09-031-485-28	US-09-188-930-334	US-08-185-432-17	US-09-113-825-2	US-08-537-210A-2	US-09-519-223-17	US-08-555-568B-17	US-09-082-059-2	US-09-065-474-38	US-08-847-429A-38	US-09-031-485-38	US-09-519-223-23	US-08-555-568B-23
18,	w.	Sequence 3, Appli	28,	28,	28,	334	17	N	2	17	17	2	38	Sequence 38, Appl	Sequence 38, Appl	Sequence 23, Appl	Sequence 23, Appl

ALIGNMENTS

US-09-439-313-376

GENERAL INFORMATION:

APPLICANT: Xu, Jiangchun APPLICANT: Dillon, Davin C. APPLICANT: Mitcham, Jennifer L. Sequence 376, Application US/09439313 Patent No. 6329505

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APPLICANT: Fanger, Gary
APPLICANT: Retter, Mark
APPLICANT: Solk, John
APPLICANT: Solk, John
APPLICANT: Day, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C9
CURRENT FILING DATE: 1999-11-12
CURRENT FALING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 376
LENGTH: 329
TYPE: PRT
ORGANISM: Homo sapien
US-09-439-313-376
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APPLICANT:
121
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                                                                                                                                                                                                  HFWRPQRLLCEDAWEQEVQVVLFLLPLLQGSGKSNVVAWGDYDDSAFMDPRYHVHGEDLD 120
                                                                    KLHRAAWWGKVPRKDLIVMLRDTDVNKRDKQKRTALHLASANGNSEVVKLVLDRRCQLNV
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Reed, Steven G
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Seque...
Patent No. 632700.

GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.

The locker, Susan Louise
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C9
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 532, Appresent No. 632950
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Best Local
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APPLICANT:
                                                                                 Sequence 379, Application US/09439313 Patent No. 6329505
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APPLICANT: Dillon, Davi
APPLICANT: Mitcham, Jen
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TYPE: PRT
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Local Similarity 98.8%;
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Retter, Mark
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Reed, Steven G
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Mitcham, Jennifer L.
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Pred. No. 2.7e-135;
1; Mismatches 2;
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; ORGANISM: Homo sapien US-09-439-313-380
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CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Kalos, Micha APPLICANT: Fanger, Gary APPLICANT: Retter, Mark APPLICANT: Solk, John APPLICANT: Day, Craig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                               NUMBER OF SEQ ID NOS: 575
SOFTWARE: FASTSEQ for Windows Version
SEQ ID NO 380
LENGTH: 671
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                                                                                                                 APPLICANT: SOIK, JOHN
APPLICANT: Day, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121-42709
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 1999-11-12
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                                                                                                                                                                                                                                                                                                                                        APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer L.
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EHGTDPNIPDEYGNTTLHYAVYNEDKLMAKALLLYGADIESKNKHGLTPLLLGIHEQKQQ 265
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NKKDKQKRTALHLASANGNSEVVKLLLDRRCQLNVLDNKKRTALIKAVQCQEDECALMLL
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Reed, Steven G
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Retter, Mark
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Reed, Steven G
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Retter, Mark
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93.9%;
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Pred. No. 2.5e-121;
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Query Match

Score 1136;

DB 4;

Best

Similarity

93.9%;

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RESULT 6
US-09-439-313-377
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER FILE REFERENCE: 210121.427C9
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 378
LENGTH: 1719
                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
Matches 216; Conserv
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Best Local
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APPLICANT:
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No. 6329505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 378,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Xu, Jiangchun APPLICANT: Dillon, Davi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                 266 VVKFLIKKKANLNALDRYGRTALILAVCCGSASIVSPLLEONVDVSSQDL 315
                                                                                                                                                                                                                          167 NKKDKQKRTALHLASANGNSEVVKLLLDRRCQLNVLDNKKRTALIKAVQCQEDECALMLL 226
                                                                                                                                                                                                                                                                          146
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-439-313-378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              YPE: PRT
                                                                                          287
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                                                                                                                                                                                                                                                                                                                    86 PLLQGSGKSNVVAWGDYDDSAFMDPRYHVHGEDLDKLHRAAWWGKVPRKDLIVMLRDTDV 145
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   86
                                                                                VVKFLIKKKANLNALDRYGRTALILAVCCGSASIVSLLLEQNIDVSSQDL 336
                                                                                                                                                                                                                                                 NKRDKQKRTALHLASANGNSEVYKLYLDRRCQLNVLDNKKRTALTKAVQCQEDECALMLL 205
                                                                                                                                                                                                                                                                                                 PCCRGSGKSKVGAWGDYDDSAFMEPRYHVRGEDLDKLHRAAWWGKVPRKDLIVMLRDTDV 166
                                                                                                                                                     EHGTDPNIPDEYGNTTLHYAVYNEDKLMAKALLLYGADIESKNKHGLTPLLLGIHEGKQQ 265
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        VVKFLIKKKANLNALDRYGRTALILAVCCGSASIVSDLLEQNVDVSSQDL 315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EHGTDPNIPDEYGNTTLHYAIYNEDKLMAKALLLYGADIESKNKHGLTPLLLGVHEQKQQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NKKDKQKRTALHLASANGNSEVVKLLLDRRCQLNVLDNKKRTALIKAVQCQEDECALMLL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NKRDKQKRTALHLASANGNSEVVKLVLDRRCQLNVLDNKKRTALTKAVQCQEDECALMLL 205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCCRGSGKSKVGAWGDYDDSAFMEPRYHVRGEDLDKLHRAAWWGKVPRKDLIVMLRDTDV 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PLLQGSGKSNVVAWGDYDDSAFMDPRYHVHGEDLDKLHRAAWWGKVPRKDLIVMLRDTDV 145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      VVKFLIKKKANLNALDRYGRTALILAVCCGSASIVSLLLEQNIDVSSQDL 336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fanger, Gary
Retter, Mark
Solk, John
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Kalos, Michael
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Day, Craig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Harlocker, Susan Louise
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Mitcham, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/09439313
                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                               66.3%;
                                                                                                                                                                                                                                                                                                                                                                            Score 1136; DB 4; Length 1719;
Pred. No. 1.2e-120;
7; Mismatches 7; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No. 2.6e-121;
7; Mismatches 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                            0;
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                                                                                                                                                       286
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APPLICANT: SOIK, John
APPLICANT: DAY, CRA19

APPLICANT: DAY, CRA19

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.42709

CURRENT APPLICATION NUMBER: US/09/439,313

CURRENT APPLICATION NUMBER: US/09/439,313

CURRENT FILING DATE: 1999-11-12

NUMBER OF SEQ ID NOS: 575

SOFTWARE: FASTSEQ for Windows Version 3.0

SEQ ID NO 377

LENGTH: 148

TYPE: PRT
ORGANISM: Homo sapien
FEATURE:
FEATURE:
NAME/KEY: VARIANT
LOCATION: (1)...(148)
OTHER INFORMATION: Xaa = Any Amino Acid

US-09-439-313-377
                                                                                                                                                                                                                                                                                                                                                                  US-09-031-485-33; Sequence 33, Appli.; Patent No. 5824306
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APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin
APPLICANT: Mitcham, Jenn!
APPLICANT: Harlocker, Su
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Best Local Similarity 96.7%;
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                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 377, Application US/09439313 Patent No. 6329505
                                                                                                                                                                                                                TITLE OF INVENTION: DIRC
TITLE OF INVENTION: PROJ
TITLE OF INVENTION: USE:
NUMBER OF SEQUENCES: 85
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 9:
                                                                                                                                                                                                                                                                                                          APPLICANT: Tang, Liang
APPLICANT: Blehm, E. Scot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   127 WWGKVPRKDLIVMLRDTDVNKRDKOKRTALHLASANGNSEVVKLVLDRRCQLNVLDNKKR 186
                                                                                                                              STREET: 1825 Sharp
CITY: Fort Collins
STATE: Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 145 KNK 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    247 KNK 249
                                                                                           COUNTRY: U
       SOFTWARE:
                                                                                                                                                                                    ADDRESSEE: Carol ADDRESSEE: Heska
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TALTKAVQCQEDECALMLLEHGTDPNIPDEYGNTTLHYAVYNEDKIMAKALLLYGADIES 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WWGKVPRKDLIVMLRDTDVNKXDKQKRTALHLASANGNSEVVKLXLDRRCQLNVLDNKKR 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fanger, Gary
Retter, Mark
Solk, John
                                                                                                                                                                                                                                                                                                                                                                                   Application US/09031485
                                                                                                                                                           1825 Sharp Point Drive
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Harlocker, Susan Louise
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                                                                                                               USA
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SYSTEM: Windows 95
WordPerfect for Windows, Version
                                                                                                                                                                                Heska Corporation
                                                                                                                                                                                                 Talkington Verser,
                                                                                                                                                                                                                                                       DIROFILARIA AND BRUGIA ANKYRIN PROTEINS, NUCLEIC ACID MOLECULES, USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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Pred. No. 1.2e-63;
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4; Indels

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Gaps

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; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-09-031-485-33
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                                                                                                                                                                                                                                                                                                          Sequence 33, Patent No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 970/484-9505
INFORMATION FOR SEQ ID NO: 33:
                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Verser, Carol Talk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 1745 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: 970/493-7272
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA: US APPLICATION NUMBER: US
                                                                                                                                                                                              APPLICANT: Tang, Liang
APPLICANT: Blehm, E. Scot
TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
TITLE OF INVENTION: PROTEINS, NUCLEIC ACID MOLECULES,
TITLE OF INVENTION: USES THEREOF
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                            NUMBER OF SEQUENCES: E
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        394 KIVELLLKYHAAIEATTESGLSPLHVAAFMGAINIVIYLLQQGANADVA-----TVRGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: 9/0/12-
TELEPHONE: 9/0/484-9505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/0 FILING DATE: 24-APR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                623 QMDIASTLLHYKANANAESKAGFTPLHLAAQEGHREMAALLIENGAKVGAQ 673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          236 ALLLYG--ADIESKN-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Verser, Carol Talkington REGISTRATION NUMBER: 37,459 REFERENCE/DOCKET NUMBER: HW-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        76 QEVQVVLP-----LLPLLQGS--GKSNVVAW----GDYDDSAFMDPRYHVHGE 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17 LHLAGS-DILSRSLMAEEYTIVHASFISCISSSLDGQGERQEQRGHFWRPQRLLCEDAWE 75
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                                                    COUNTRY:
                                                                     CITY: Fort Collins
STATE: Colorado
                                                                                                      ADDRESSEE: Carol Talkington Verser, Ph.D. ADDRESSEE: Heska Corporation STREET: 1825 Sharp Point Drive
                                     ZIP:
                                                                                                                                                                                                                                                                                                                                                                                                                                         KQQVVKFLIKKKANLNALDRYGRTALILAVCCGSASIVSPLLEQNVDVSSQ 313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ASPNAATRDLYTPLHIAAKEGQEEVAAILMDHGTDKTLLTKKGFTPLHLAAKYGNLPVAK 562
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COLNVLDNKKRTALTKAVQCQEDECALMILEHGTDPNIPDEYGNTTLHYAVYNEDKLMAK 235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  --TPLHLAA---RANQTDIVRVLVRNGAQVDAAARELQTPLHIASRLGNTDIVILLLQAN 502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DLDKLHRAAWWGKVPRKDLI-VMLRD-TDVNKRDKQKRTALHLASANGNSEVVKLVLDRR 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SLLERGTPVDIEGKNOVTPLHVAAHYNNDKVALLLLENGASAHAAAKNGYTPLHIAAKKN 622
                                80525
                                                                                                                                                                                                                                                                                                          3, Application US/08847429A 5827692
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                                                        USA
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US-08-847-429A-33
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                                                                                                                                                                                                                                                                                                                                                                              Sequence 33, Application US/09065474 Patent No. 6063599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 970/484-9505
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: WordPerfect for Windows, CURRENT APPLICATION DATA:
APPLICATION NUMBERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: HW TELECOMMUNICATION INFORMATION: 970/493-7272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Verser, Carol Talkington
REGISTRATION NUMBER: 37,459
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 1745 amino acids
                                                                                                                                                                                                APPLICANT: Tang, Liang
APPLICANT: Blehm, E. Scot
TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
TITLE OF INVENTION: PROTEINS, NUCLEIC ACID MOLECULES,
TITLE OF INVENTION: USES THEREOF
NUMBER OF SEQUENCES: 171
CORRESPONDENCE ADDRESS:
                                                                    STREET: 1822 CITY: Fort Collins
STATE: Colorado
COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            394 KIVELLLKYHAAIEATTESGLSPLHVAAFMGAINIVIYLLQQGANADVA-----TVRGE 447
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   334 LHMAAQVDDVTVDYLTPLHVAAHCGHVRVAKLLLDRNADPNARALNGFTPLHIACKKNRI 393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 263 KQQVVKFLIKKKANLNALDRYGRTALILAVCCGSASIVSPLLEQNVDVSSQ 313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              503 ASPNAATRDLYTPLHIAAKEGQEEVAAILMDHGTDKTLLTKKGFTPLHLAAKYGNLPVAK 562
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               176
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    623 QMDIASTLLHYKANANAESKAGFTPLHLAAQEGHREMAALLIENGAKVGAQ 673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      236 ALLLYG--ADIESKN-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
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                                                                 COUNTRY:
                                                                                                                                                        ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QEVQVVLP------LLPLLQGS--GKSNVVAW----GDYDDSAFMDPRYHVHGE 117
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SLLERGTPVDIEGKNOVTPLHVAAHYNNDKVALLLLENGASAHAAAKNGYTPLHIAAKKN
                                            80525
                                                                                                                                   1825 Sharp Point Drive
                                                                                                                                                             Heska Corporation
                                                                                                                                                                                  Carol Talkington Verser,
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  Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -----KHGLTPLLLGIHEQ 262
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-08-810-712-10
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                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Yeda Research and Development Co. LTD
TITLE OF INVENTION: Tumor Suppressor Genes, Proteir
TITLE OF INVENTION: Use of said Genes and Proteins
FILE REFERENCE: sequencelist
CURRENT APPLICATION NUMBER: US/08/810,712G
CURRENT FILING DATE: 1997-03-03
BARLIER APPLICATION NUMBER: PCT/US94/11598
EARLIER FILING DATE: 1994-10-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-065-474-33
                                                                                     SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                      Sequence 10, Application US/08810712G Patent No. 6160106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
                                                                                                                               NUMBER OF SEQ ID NOS: 31
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INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
                                                                   LENGTH: 1423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Verser, Carol Talkington REGISTRATION NUMBER: 37,459
REFERENCE/DOCKET NUMBER: HW-5-C
TELECOMMUNICATION INFORMATION:
TELEPHONE: 970/493-7272
TELEFAX: 970/484-9505
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SOFTWARE: WordPerfect for Windows,
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            563 SILERGTPVDIEGKNQVTPLHVAAHYNNDKVALLLLENGASAHAAAKNGYTPLHIAAKKN 622
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 118 DLDKLHRAAWWGKVPRKDLI-VMLRD-TDVNKRDKQKRTALHLASANGNSEVVKLVLDRR 175
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    236 ALLLYG--ADIESKN----
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TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17 LHLAGS-DLLSRSLMAEEYTIVHASFISCISSSLDGQGERQEQRGHFWRPQRLLCEDAWE 75
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Local Similarity 25.1%; Pred. No. 5.8e-15;
les 88; Conservative 57; Mismatches 141
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       --TPLHLAA---RANQTDIVRVLVRNGAQVDAAARELQTPLHIASRLGNTDIVILLLQAN 502
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Query Match
Best Local Similarity
"~+~hes 68; Conserv
                                                                                                                     ; TOPOLOGY: linear; MOLECULE TYPE: Protein US-09-065-474-139
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Patent No. 60
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Best Local Similarity
Matches 71; Conserva
                                                                                                                                                                                                                                  TELEFAX: 970/484-9505
INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Verser Carol Talkington
REGISTRATION NUMBER: 37,459
REFERENCE/DOCKET NUMBER: HW-5-C
TELECOMMUNICATION INFORMATION:
TELEPHONE: 970/493-7272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 80525
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 352 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: WordPerfect for Windows,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Tang, Liang
APPLICANT: Blebm, E. SCOT
APPLICANT: Blebm, E. SCOT
TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
TITLE OF INVENTION: PROTEINS, NUCLEIC ACID MOLECULES,
TITLE OF INVENTION: USES THEREOF
NUMBER OF SEQUENCES: 171
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                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: FILING DATE: 24-APPCLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Carol Talkington Verser,
ADDRESSEE: Heska Corporation
STREET: 1825 Sharp Point Drive
CITY: Fort Collins
STATE: Colorado
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IVEC---LAEHGADLNACDKDGHIALHLAVRRCQMEVIKTLLSQGCFVDYQDRHGNTPLH
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                                                                                                                                                                            amino acids
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            12.1%; Score 208; DB 3; Jilarity 28.9%; Pred. No. 2.5e-15; Conservative 38; Mismatches 89;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     USA
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                                                                                                                                                                                                                                      139:
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                                                                                                                                                                                                                                                                                                                      HW-5-C1
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                                                  Length 352;
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US-08-281-193-2
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Patent No. 5466595
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Best Local (
                                                                                                                                                                                                                                                                                                                                                         Matches
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
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                                                                          QEDECALMILEHGTDPNIPDEYGNTTLHYAVYNEDKLMAKALLLYGADIESKNKHGLTPL 255
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                                                       LLGIHEQKQQVVKFLIKKKANLNALDRYGRTALILA 291
                                                                                                                IISMDSSQIHSKDPRYGASPLHWAK---NAEMARMLLKRGCDVDSTSAAGNTALHVAVMR
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--LQDLMPISRARKPAFILS 414
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                                                                                                                                                                                                                                                                                                                                                           Score 206.5; DB 1;
Pred. No. 1.2e-14;
3; Mismatches 117;
                                                                                                                                                                                                             ----RYHVHGEDLDKLHRAAWWGKVPRKDL
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US-08-422-106-2
; Sequence 2, Application US/08422106
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Best Local
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                                                                                                                                                                 Sequence 2, Application US/08735716 Patent No. 5840511
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APPLICANT: Jones, Simon
APPLICANT: Tang, Jim
APPLICANT: Tang, Jim
TITLE OF INVENTION: Calicium Indepen
NUMBER OF SEQUENCES: 15
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                    GENERAL INFORMATION:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/422,10
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PRIOR APPLICATION DATA:
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22.9%; Pred. No. 1.2e-14;
tive 53; Mismatches 117
                                                                                               Calcium Independent Phospholipase A2/B
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LENGTH: 752 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-735-716-2
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PRIOR APPLICATION DATA:
APPLICATION UNMBER: US/08/281,193
FILING DATE: 27-JUL-1994
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 752 amino acids
                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
ATTORNEY/AGENT INFORMATION: NAME: Brown, Scott A.
                                                                                                                                                                                                                                                                                   APPLICANT: Jones, Simon
APPLICANT: Tang, Jim
TITLE OF INVENTION: Calcium Independent Phospholipase A2/B
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute, Inc.
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APPLICATION NUMBER: US,
FILING DATE: 23-OCT-19
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                                                      APPLICATION NUMBER: FILING DATE:
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                                        CLASSIFICATION:
                                                                                                                                                                                                                                                      STREET: 87 Cambridge
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87 CambridgePark Drive
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                                                                   US/08/555,568B
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; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-555-568B-2
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
TELEPHONE: (617) 498-8224
                              256 LLGIHEQKQQVVKFLIKKKANLNALDRYGRTALILA 291
                                                                          328
                                                                                              196 QEDECALMILEHGTDPNIPDEYGNTTLHYAVYNEDKLMAKALLLYGADIESKNKHGLTPL
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   388 FMASKISKQ-
                                                                                                                                                                                                                                                                    158 LACRKGDSEILVELVQYCHAQMDVTDNKGETAFHYAVQGDNSQVLQLLGKNASAGLNQVN 217
                                                                                                                                                                                                                                   98 AWGDYDDSAFMDP------136
                                                                                                                                                                                                                                                                                                                   68
                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 22.9%; nes 77; Conservation
                                                                                                                                                                                                                                                                                                                                                           8 SHPLWVDSFLHLAGSDLLSRSLMAEEYTIVHASFISCISSSLDGOGERQEQRGHFWRPQR 67
                                                             NRFDCVMVLLTYGANAGTPGEHGNTPLHLAISKDNMEMIKALIVFGAEVDTPNDFGETPA
                                                                                                                                                                                                                                                                                                                   LLC-----
---LQDLMPISRARKPAFILS 414
                                                                                                                                                                                                                                                                                                       -----EDAWEQEVQ----VVLPLLPLLQGSGKSNVV 97
                                                                                                                                                                                                                                                                                                                                                                                                                53; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                              Score 206.5; DB 2;
Pred. No. 1.2e-14;
53; Mismatches 117;
                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 752;
                                                                                                                                                                                                                                                                                                                                                                                                              89;
                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                 387
                                                                                                                                                                                                      270
```

10;

Search completed: August 2, 2002, 08:44:06 Job time: 276 sec

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Page 8





ALIGNMENTS

```
RESULT 1
US-09-439-313-376
; Sequence 376, Application US/09439313
; Patent No. 6329505
; GENERAL INFORMATION:
; APPLICANT: Xu, Jiangchun
; APPLICANT: Dillon, Davin C.
; APPLICANT: Mitcham, Jennifer L.
; APPLICANT: Harlocker, Susan Louise
; APPLICANT: Harlocker, Susan Louise
; APPLICANT: Reed, Steven G.
; APPLICANT: Kalos, Michael
; APPLICANT: Fanger, Gary
; APPLICANT: Fetter, Mark
; APPLICANT: Solk, John
; APPLICANT: Oay, Craig
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
; TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
; FILE REFERENCE: 210121.427C9
; CURRENT APPLICATION NUMBER: US/09/439,313
; CURRENT FILING DATE: 1999-11-12
; NUMBER OF SEQ ID NOS: 575
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 376
; LENGTH: 329
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-439-313-376
```

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Query Match 100.0%; Score 1713; DB 4; Length 329; Best Local Similarity 100.0%; Pred. No. 5.7e-188; Matches 329; Conservative 0; Mismatches 0; Indels 0
                                                  0; Gaps
     Qy
Db
     61 HFWRPQRLLCEDAWEQEVQVVLPLLPLLQGSGKSNVVAWGDYDDSAFMDPRYHVHGEDLD 120
Qy
     61 HFWRPQRLLCEDAWEQEVQVVLPLLPLLQGSGKSNVVAWGDYDDSAFMDPRYHVHGEDLD 120
Db
    121 KLHRAAWWGKVPRKDLIVMLRDTDVNKRDKQKRTALHLASANGNSEVVKLVLDRRCQLNV 180
Qy
    Db
    Οv
Db
```

```
        Qy
        241 GADIESKNKHGLTPLLLGIHEQKQQVVKFLIKKKANLNALDRYGRTALILAVCCGSASIV
        300

        Db
        241 GADIESKNKHGLTPLLLGIHEQKQQVVKFLIKKKANLNALDRYGRTALILAVCCGSASIV
        300

        Qy
        301 SPLLEQNVDVSSQDLERRPESMLFLVIIM
        329

        Db
        301 SPLLEQNVDVSSQDLERRPESMLFLVIIM
        329
```

RESULT 2

